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Microbial utilization of residue carbon depends on placement and differs in humid and semi-arid climates

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Canada 

Introduction

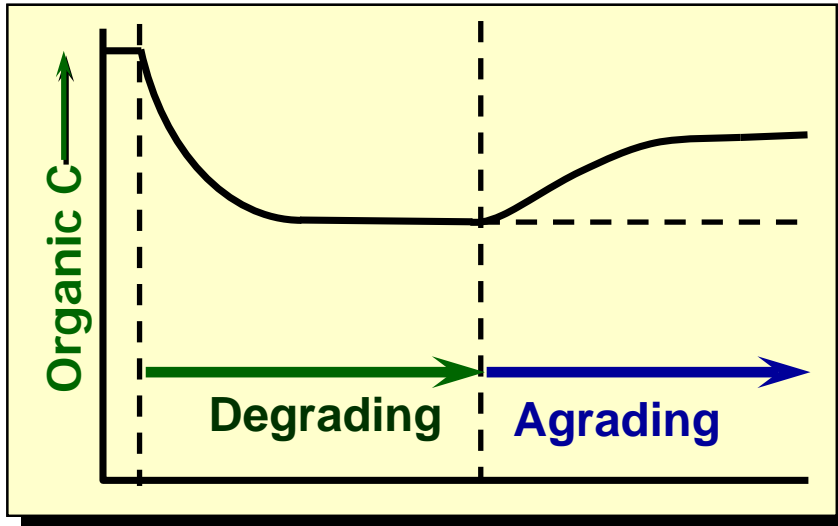
- Producing harvestable C is the main goal of agriculture
- Crop residues are the major source of new C in agricultural soils

Sustain long-term soil productivity

- rates and mechanisms of residue C decomposition
- microbial regulation



Introduction



$$\text{C storage} = \text{C input} - \text{decomposition}$$

H. Janzen

- By manipulating residue placement we can alter the rate and perhaps pathway of C decomposition

Introduction

Microbes are primary regulators of litter decay, but pathways of decomposition are only vaguely understood and poorly quantified



Breton: triticale



Swift Current: lentil

Use of ^{13}C *enriched plant* material allows us to track fate of C in various physical, chemical and biological pools

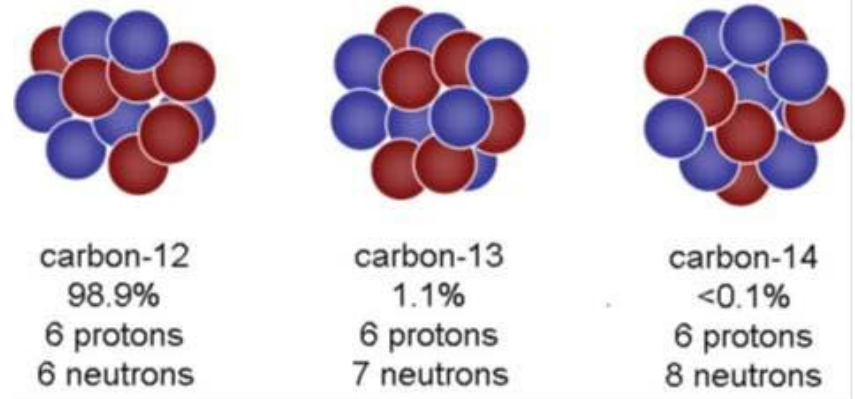
“stable isotope probing”

Background: Stable Isotope Probing

- What is a stable isotope?

Common carbon isotopes: ^{12}C , ^{13}C , ^{14}C
6 protons (6, 7 and 8 neutrons)

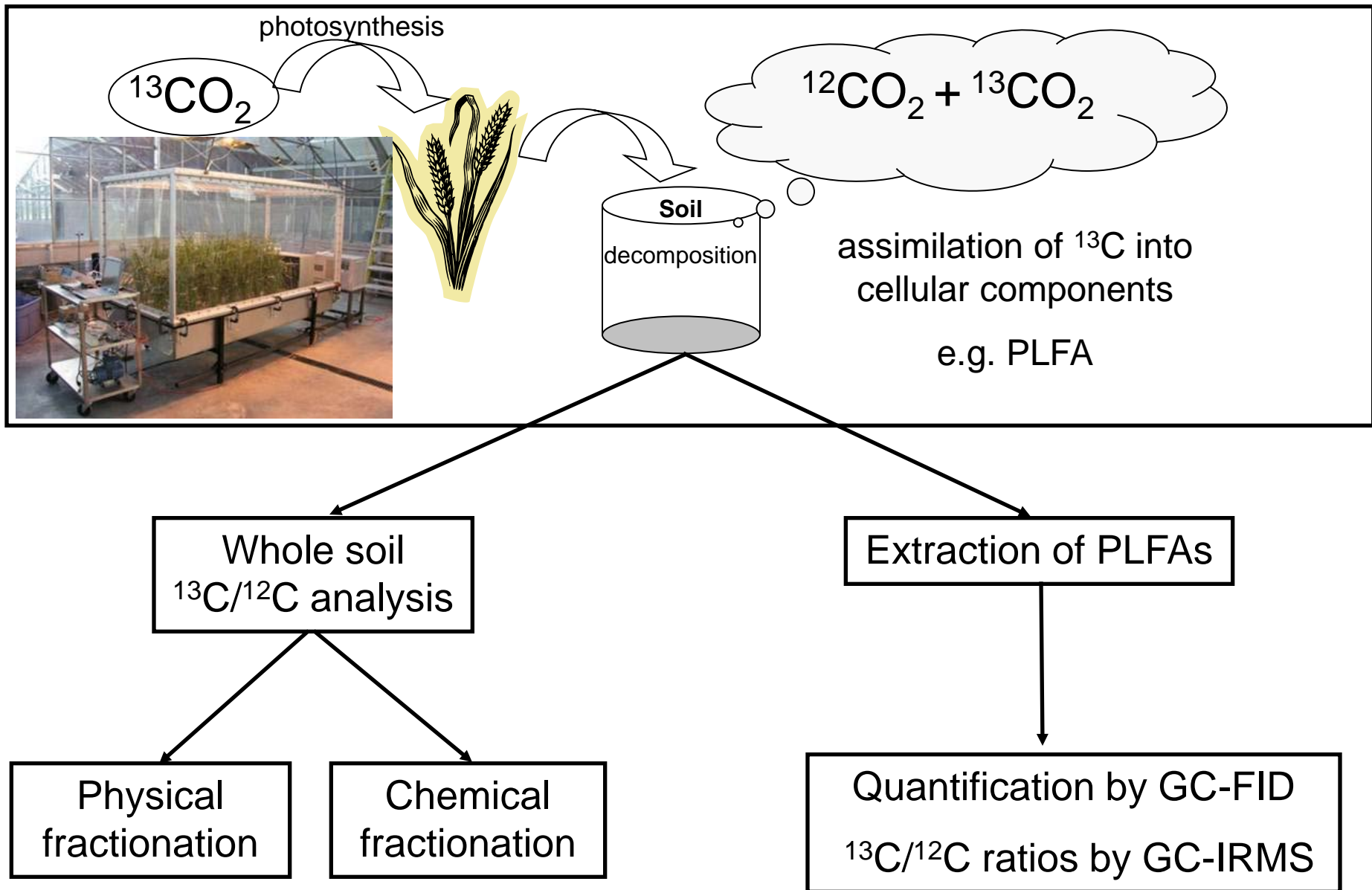
- ^{12}C and ^{13}C are stable (non-radioactive) and occur naturally at a ratio of ca. 99:1



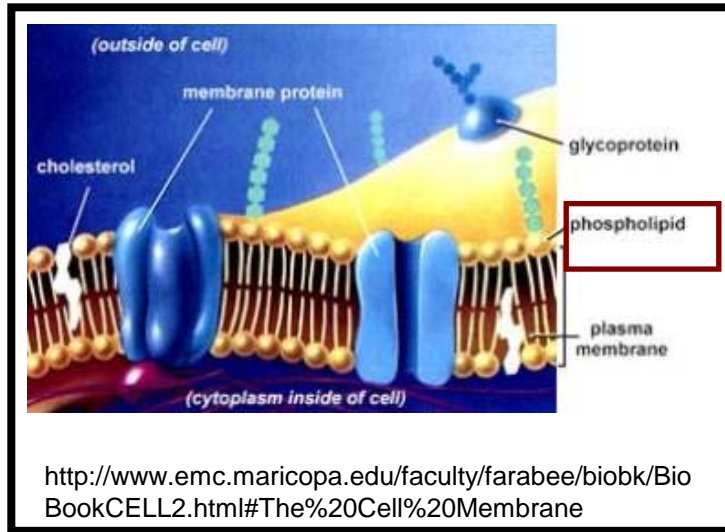
<http://wordpress.mrreid.org/2011/03/23/>

- For our purposes ^{13}C acts as a *tracer*.

Background: Stable Isotope Probing



SIP/Phospholipid fatty acid analysis



Active populations:

1. abundance
2. microbial community structure

Microorganisms metabolize C obtained from their environment (^{12}C + ^{13}C)

Coupled with ^{13}C -SIP, PLFA indicates *functionally active* components of the soil microbial community



**Microbial utilization of residue carbon
depends on placement and differs in humid
and semi-arid climates**

Objective:

**to determine the effect of residue placement on
retention of ^{13}C in the microbial biomass in humid
and semi-arid climates**

Experimental Design

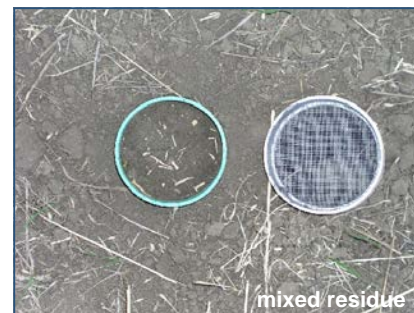
Fall 2007, microcosms set up in transects

Each transect (n=4) has 10 microcosms



Barley residue (^{13}C at ca. 10 atom %) added at a rate of 2 Mg C ha^{-1} , either:

- a. incorporated (0-10cm)
- b. surface applied



Experimental design

**Microcosms are amended annually in fall with *non-labeled* barley residue



^{13}C residue added

Microcosm destructive sampling

20 yr?

Fall
2007

Spring
2008

Fall
2008

Fall
2009

Fall
2010

Microcosm sampling

- Microcosms (n=4) destructively sampled:
 - 6 mo (spring 2008)
 - 12 mo (fall 2008)
 - 24 mo (fall 2009)



- Cores divided into 2 depths:
 - 0-5cm
 - 5-10cm
- PLFA analysis:
 - GC-FID, MIDI (quantitative)
 - GC-IRMS ($^{13}\text{C}/^{12}\text{C}$ of 26 individual PLFAs)

Experimental design:

Basic Site characteristics:

Ottawa: humid

fine sandy loam

MAP: 882mm; MAT: 5.8°C



Lethbridge: semi-arid

sandy clay loam

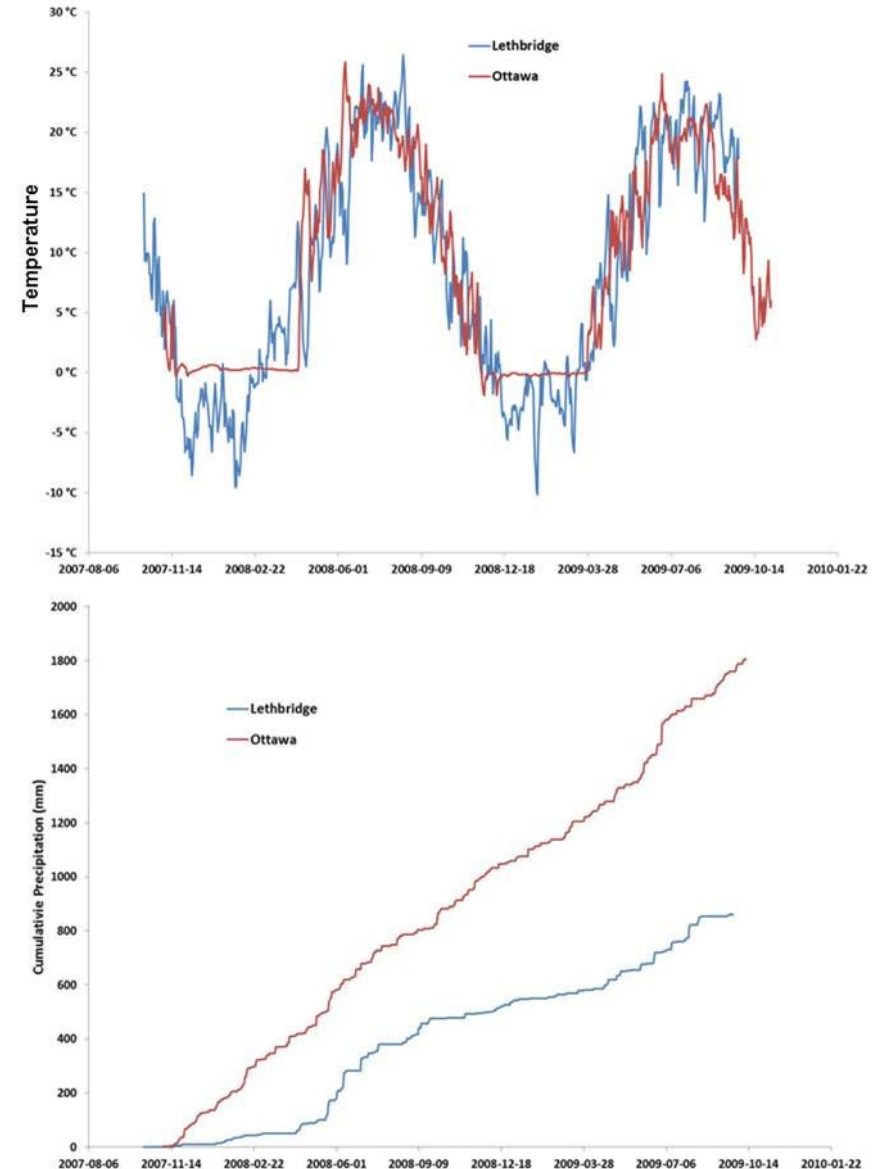
MAP: 386mm; MAT: 5.7°C



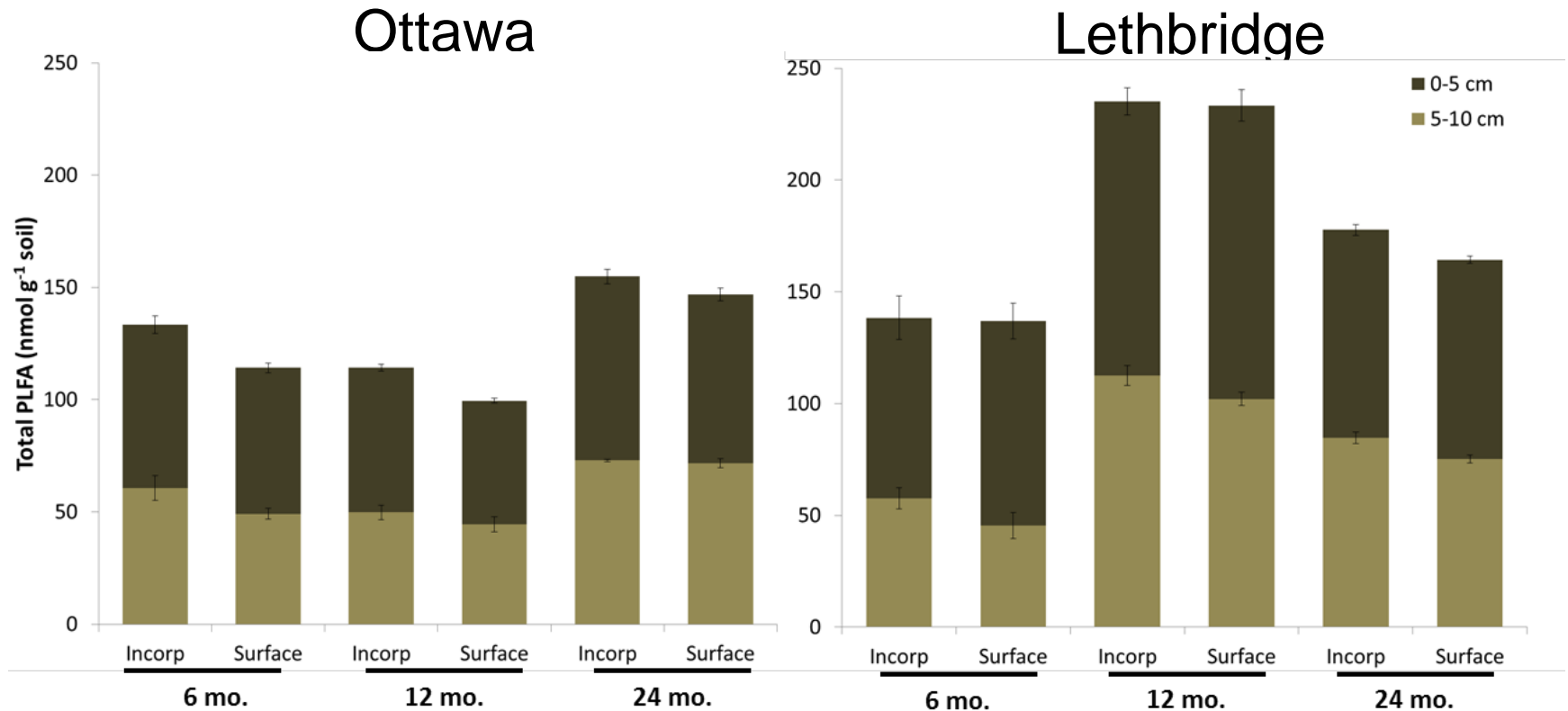
Experimental design

Basic Site characteristics:

- similar mean annual temperature
- very different temporal patterns of soil temperature
- very different precipitation patterns



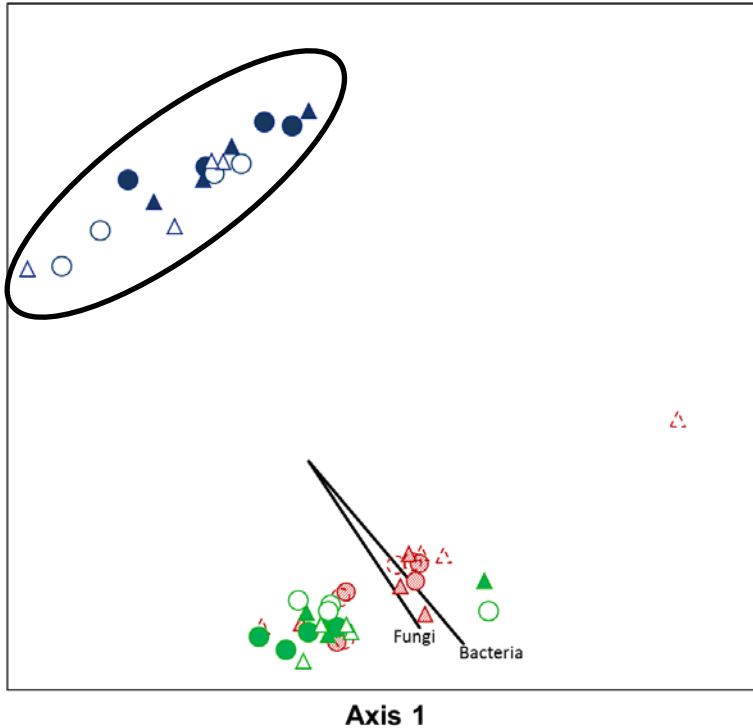
Microbial abundance – ^{12}C + ^{13}C PLFA



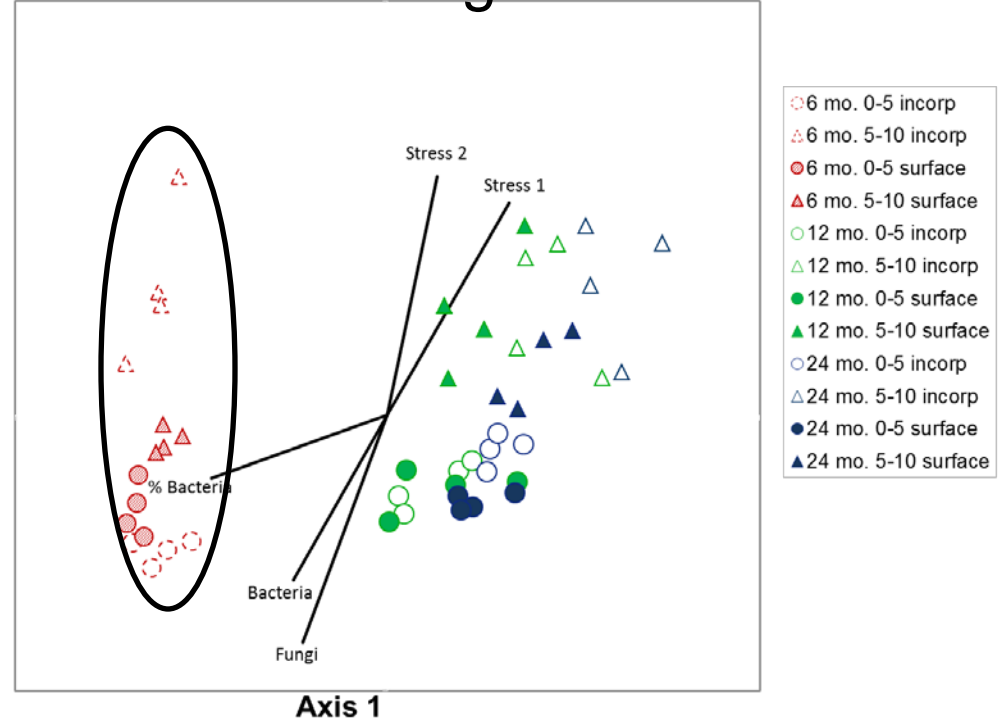
- Higher fall biomass at Lethbridge than Ottawa
- Significant biomass at 0-5 and 5-10 cm depths

Community structure – ^{12}C + ^{13}C PLFA

Ottawa



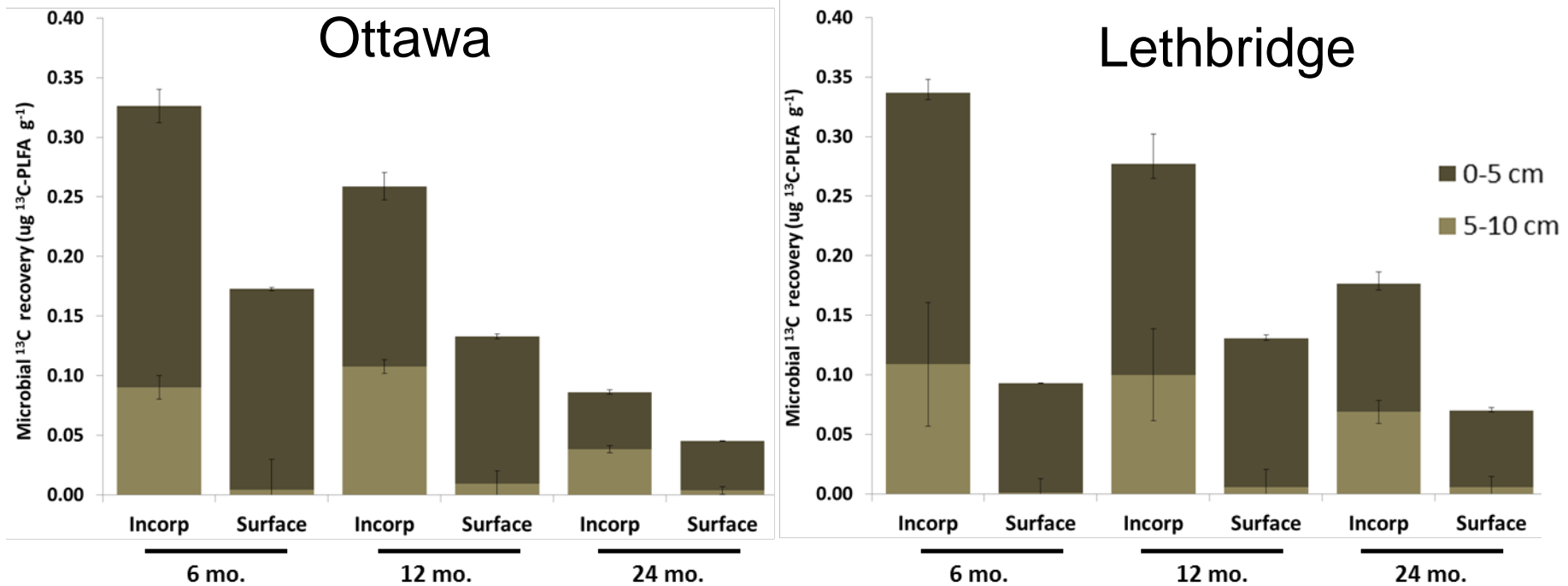
Lethbridge



Ottawa: communities similar in season 1, major shift by 24 mo.

Lethbridge: strongest effects of season and depth

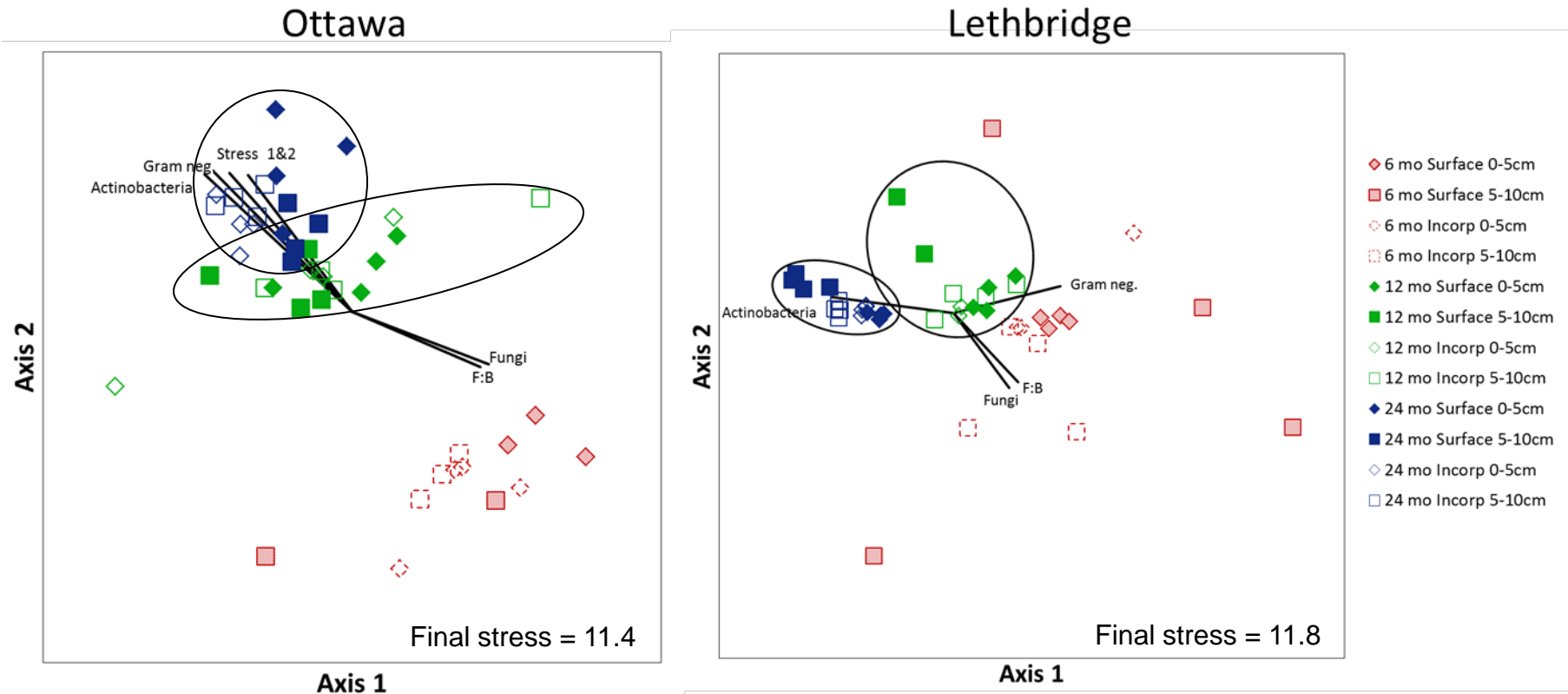
^{13}C recovery in the microbial biomass



Incorporated: Highest recovery at 6 mo.
Little difference between sites

Surface applied: Only 50% as much ^{13}C recovered
Negligible ^{13}C recovery below 5 cm after 2 yr

^{13}C recovery in the microbial biomass



- Initially, ^{13}C labelled community structure highly variable
- Stronger effect of depth at Lethbridge than at Ottawa

^{13}C PLFA Community structure: 24 mo.



2 years after the application of the ^{13}C label:

- distinct community structure reflected earlier residue placement
- there was a strong effect of depth at Lethbridge

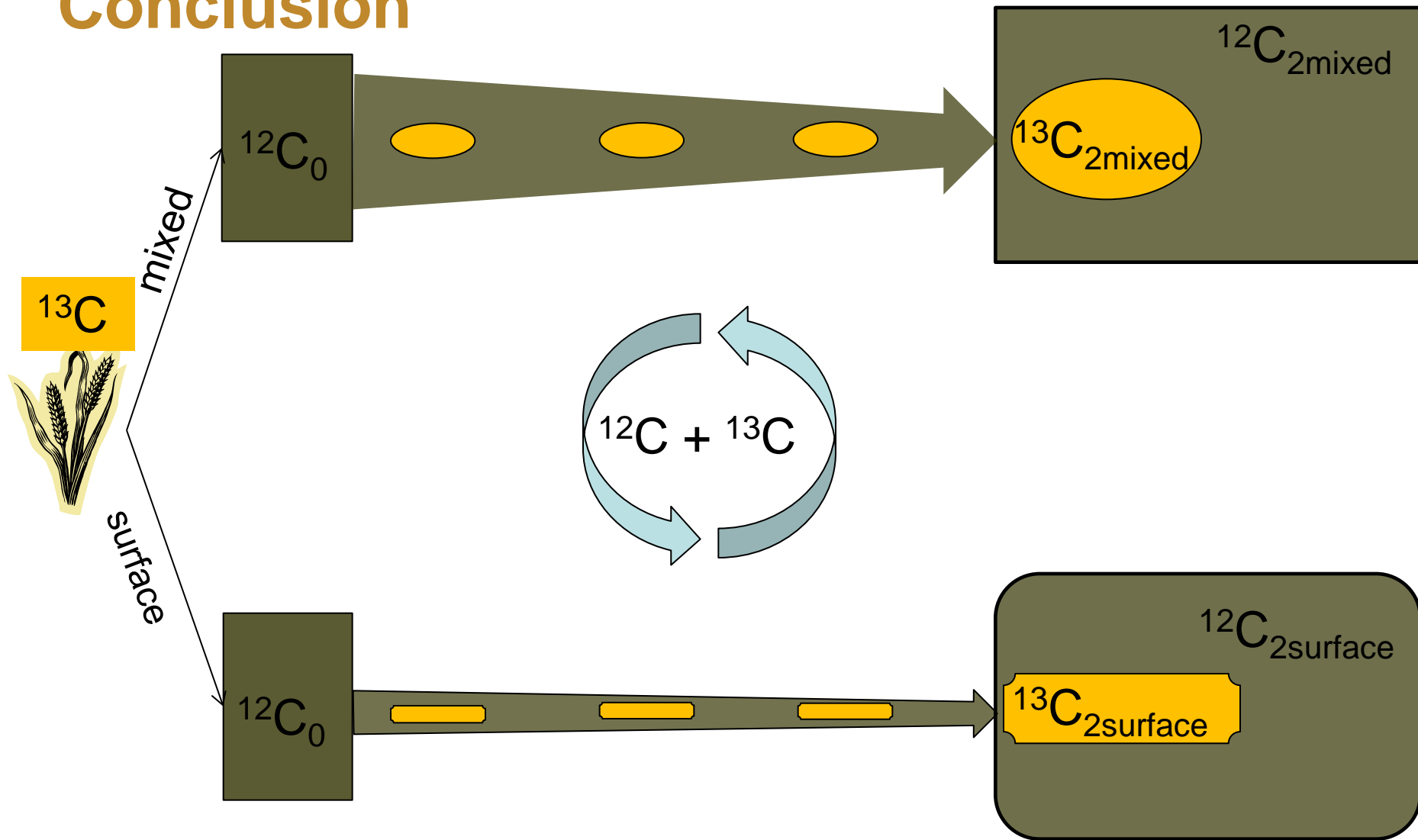
Conclusion

The effect of residue placement was:

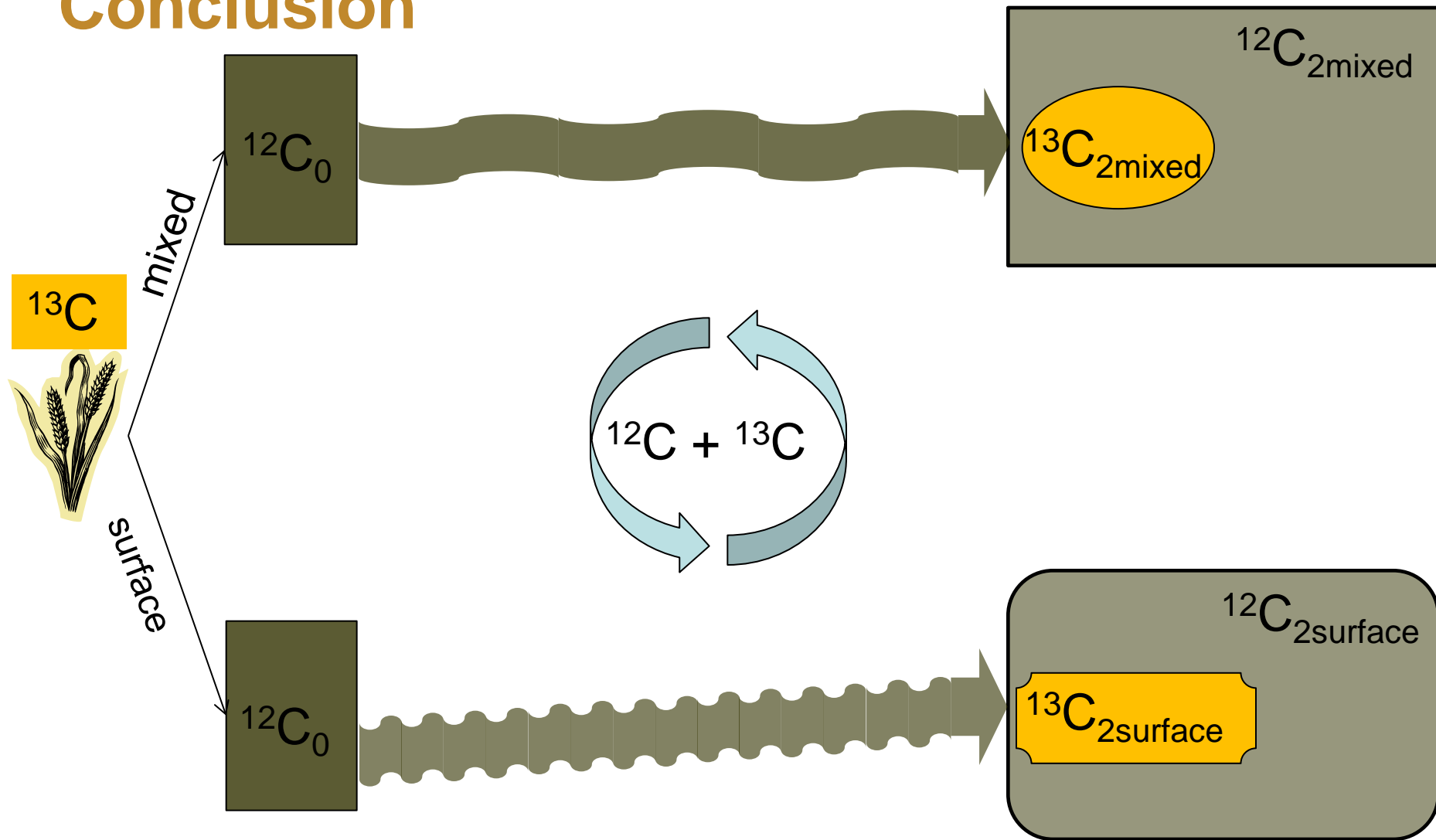
- *site-specific* with,
- a gradual shift toward a *defined community structure* of the labeled microbial biomass that reflected whether residues had been incorporated into the soil or applied to the surface *two years earlier*.



Conclusion



Conclusion



Implication

Climate has a greater influence on microbial utilization of residues when they are placed on the surface than when they are mixed into the soil.

- Consequently, residue C dynamics may show much higher-site-to-site variability in no-tillage systems than in conventionally tilled systems
- This may explain why NT increases soil C in some cases, but not in others

Acknowledgements

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